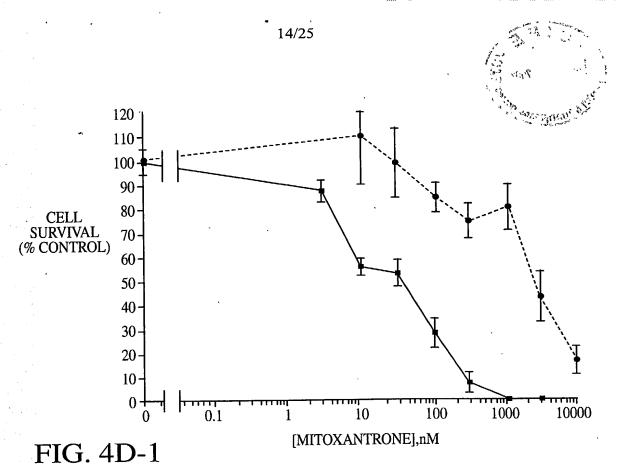
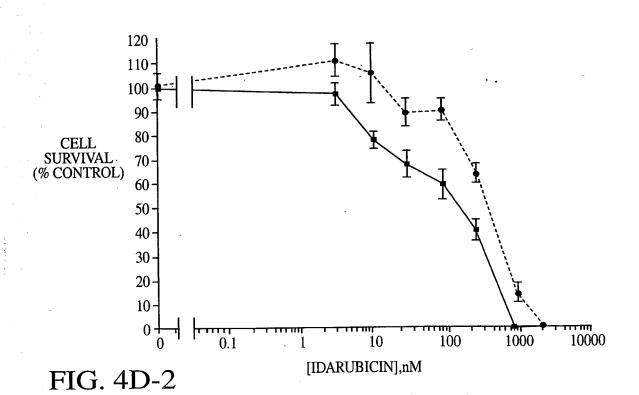


| 901 | AAATGCTGTC CTTTTGCTCC TGAAAAGGAT GTCTAAGCAG GGACGAACAA |
|------|--|
| 951 | TCATCTTCTC CATTCATCAG CCTCGATATT CCATCTTCAA GTTCTTTGAT |
| 1001 | AGCCTCACCT TATTGGCCTC AGGAAGACTT ATGTTCCACG GGCCTGCTCA |
| • | GGAGGCCTTG GGATACTTTG AATCAGCTGG TTATCACTGT GAGGCCTATA |
| 1051 | ATAACCCTGC AGACTTCTTC TTGGACATCA TTAATGGAGA TTCCACTGCT |
| 1101 | GTGGCATTAA ACAGAGAAGA AGACTTTAAA GCCACAGAGA TCATAGAGCC |
| 1151 | TTCCAAGCAG GATAAGCCAC TCATAGAAAA ATTAGCGGAG ATTTATGTCA |
| 1201 | TTCCAAGCAG CMTTACATCA ACTTTCCGGG |
| 1251 | ACTCCTCCTT CTACAAAGAG ACAAAAGCIG AATTMOMEDA GCTACAAGACACGAC GGTGAGAAGA AGAAGAAGAT CACGGTCTTC AAGGAGATCA GCTACACGAC |
| 1301 | \mathcal{L} |
| 1351 | CTCCTTCTGT CATCAACTCA GITTOUT ACARCATTCT CACAGTCGTA |
| 1401 | TGCTGGGTAA TCCCCAGGCC TGTTTTTT |
| 1451 | CTGGGACTGG TTATAGGTGC CATTTACTTT GGGCTAAAAA ATGATTCTAC |
| 1501 | TGGAATCCAG AACAGAGCTG GGGTTCTCTT CTTCCTGACG ACCAACCAGT |
| 1551 | GTTTCAGCAG TGTTTCAGCC GTGGAACTCT TTGTGGTAGA GAAGAAGCTC |
| 1601 | TTCATACATG AATACATCAG CGGATACTAC AGAGTGTCAT CTTATTTCCT |
| 1651 | TGGAAAACTG TTATCTGATT TATTACCCAT GACGATGTTA CCAAGTATTA |
| 1701 | TATTTAÇCTG TATAGTGTAC TTCATGTTAG GATTGAAGCC AAAGGCAGAT 5' PCR PRIMER (SENSE) |
| | GCCTTCTTCG TTATGATGTT TACCCTTATG ATGGTGGCTT ATTCAGCCAG |
| 1751 | TTCCATGGCA CTGGCCATAG CAGCAGGTCA GAGTGTGGTT TCTGTAGCAA |

FIG. 2C-2





0.4



PACLITAXE RF. CIAPLATIN 4,700 3,700 IDARUBICIN 140 285 DEXPRUBICIN RF **0598 LC50 DAUNORUBICIN 35.5 RF 1667** LC50 3333 MITOZANTRONS RF **000'08 1,495** 393** LC50 MCF-7/BCRPc19 MCF-7/BCRPcS MCF-7/pcDNA3 MCF-7/BCRPall MCF-7/AdrVp MCF-7/W

** = DIFFERS SIGNIFICANTLY FROM MCF-7/W OR MCF-7/pcDNA3, p <0.01 (STUDENT'S t TEST) = DIFFERS SIGNIFICANTLY FROM MCF-7/W OR MCF-7/pcDNA3, p <0.05 (STUDENT'S t TEST)